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341

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International application No PCT/EP2006/010132

	FICATION OF SUBJECT MATTER C12Q1/68		
According to	o International Patent Classification (IPC) or to both national classifica	ation and IPC	
—— <u> </u>	SEARCHED		·-
	ocumentation searched (classification system followed by classification	on symbols)	
C12Q		and the second second	•
Documentat	tion searched other than minimum documentation to the extent that s	such documents are included in the fields so	earched
Electronic d	ata base consulted during the international search (name of data ba	se and, where practical, search terms used	1)
EPO-In	ternal, EMBASE, MEDLINE, PAJ, WPI Da	ata, Sequence Search	
C. DOCUMI	ENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the rele	evant passages	Relevant to claim No.
A	NAKAMURA M ET AL: "DEVELOPMENT OF MICRO ARRAY FOR IDENTIFICATION OF INFECTIOUS DISEASE CAUSACTIVE BAC HUMAN"	. CTERIA IN	1-8, 10-25
	18 May 2003 (2003-05-18), ABSTRA THE GENERAL MEETING OF THE AMERIC SOCIETY FOR MICROBIOLOGY, THE SOC WASHINGTON, DC, US, PAGE(S) ABSTR XP008047725 ISSN: 1060-2011	CAN CIETY,	·
	abstract	-/	
X Furti	her documents are listed in the continuation of Box C.	X See patent family annex.	
"A" docume	ategories of cited documents : ent defining the general state of the art which is not lered to be of particular relevance tocument but published on or after the international	*T* later document published after the Inte or priority date and not in conflict with cited to understand the principle or the invention.	the application but eory underlying the
filing date cannot be considered novel or cannot be come involve an inventive step when the document which is cited to establish the publication date of another cannot be considered novel or cannot be considered no		t be considered to cument is taken alone claimed invention	
"O" docume other i	"O' document referring to an oral disclosure, use, exhibition or other means cannot be considered to involve an inventive step when the document is combined with one or more other such document is combined with one or more other such documents, such combination being obvious to a person skilled in the art		ore other such docu-
	ent published prior to the international filing date but an the priority date claimed	*&" document member of the same patent	family
	actual completion of the international search	Date of mailing of the international sea	rch report
	July 2007	26/07/2007	
Name and r	nailing address of the ISA/ European Patent Office, P.B. 5818 Patentlaan 2 NL – 2280 HV Rijswijk	Authorized officer	•
	NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016 Helliot, Bertrand		

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INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2006/010132

	ation). DOCUMENTS CONSIDERED TO BE RELEVANT	
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WANG R-F ET AL: "DNA microarray analysis of predominant human intestinal bacteria in fecal samples" August 2004 (2004-08), MOLECULAR AND CELLULAR PROBES, ACADEMIC PRESS, LONDON, GB, PAGE(S) 223-234, XP004522575 ISSN: 0890-8508 abstract; tables 1,2	1-8, 10-25
Α	LEHNER A ET AL: "Oligonucleotide microarray for identification of Enterococcus species" 1 May 2005 (2005-05-01), FEMS MICROBIOLOGY LETTERS, AMSTERDAM, NL, PAGE(S) 133-142, XP004876200 ISSN: 0378-1097 abstract	1-8, 10-25
X	EP 1 310 569 A (PRESIDENT OF GIFU UNIVERSITY) 14 May 2003 (2003-05-14)	1-6, 10-13, 15,19-25
X	WO 92/07096 A (MICROPROBE CORPORATION) 30 April 1992 (1992-04-30) page 12, paragraph 2 page 27, paragraph 2 example 6	1-6,10, 12,13
X	US 6 747 137 B1 (WEINSTOCK KEITH G [US] ET AL) 8 June 2004 (2004-06-08) column 2, lines 41-47 column 16, lines 55-60 column 19, lines 43-61 column 42, lines 5-43 table 2 claim 7	1-6,10, 12,13, 19-25
X	EP 1 344 833 A (CHIP BIOTECHNOLOGY INC DR [TW]) 17 September 2003 (2003-09-17)	1-6, 10-13, 15,16, 18-25
Y	page 2, paragraphs 8,10 page 3, paragraphs 13,18,19 examples 1,2 claim 8	7,14,17
Y	US 6 008 341 A1 (FOSTER TIMOTHY JAMES [IE] ET AL) 28 December 1999 (1999-12-28) figure 2	7,14,17

INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2006/010132

		PCT/EP2006/010132			
C(Continua	Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT				
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.			
X	WO 02/094868 A (CHIRON SPA [IT]; MASIGNANI VEGA [IT]; MORA MARIROSA [IT]; SCARSELLI MA) 28 November 2002 (2002-11-28) page 2, lines 12,13 page 2, lines 20-25 sequences 1992,3983	1-8, 10-15			
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		, i			

International application No. PCT/EP2006/010132

INTERNATIONAL SEARCH REPORT

Box II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)
This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
see additional sheet
As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
1-4 (totally), 5-8, 10-18 (partially), 19-25
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

Invention 1: claims 1-7 and 10-25 (partially)

An analytical device for direct identification and characterisation of microoragnisms in a sample or clinical specimen, wherein the device comprises species specific gene probes of at least 100 nucleotides, and in particular a device for Staphylococcus species identification, in particular for S. aureus identification, wherein, in this case, the microarray comprises the gene probe listed as SEQ ID \mbox{N}° 3 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus aureus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus aureus in a sample or clinical specimen.

Inventions 2-176: claims 1-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus aureus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° SEQ ID N° 1-2, 4-141, 790, 798, 801, 802, 808, 812, 814, 818, 825, 827, 837, 840, 843, 844, 846, 848-852, 854, 855, 859, 862, 875, 885, 896, 897, 904, 907, 908, 935, 942, 2902, 2903, and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus aureus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus aureus in a sample or clinical specimen.

Inventions 177-220: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of E. coli in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^\circ$ 142-173, 815, 833, 834, 836, 839, 857, 860, 886-887, 895, 901, 906, and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of E. coli in a sample or in a clinical specimen.

Inventions 221-258: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus epidermis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 174-208,786, 806, 826 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus epidermis in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus epidermis in a sample or clinical specimen.

Inventions 259-269: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus haemoliyticus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 209-215, 796, 803, 820, 938 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus haemoliyticus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus haemoliyticus in a sample or clinical specimen.

Inventions 270-276: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus lugdunensis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 216-221, 888 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus lugdunensis in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus lugdunensis in a sample or clinical specimen.

Inventions 277-284: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus warneri in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 224-230, 831 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus warneri in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus warneri in a sample or clinical specimen.

Inventions 285-286: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus saprophyticus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 222-223 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus saprophyticus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus saprophyticus in a sample or clinical specimen.

Inventions 287-375: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus pneumoniae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N°523-605, 793, 805, 807, 813, 858, 929 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus pneumoniae in a sample or in a clinical specimen.

A kit for the detection of Streptococcus pneumoniae in a sample or clinical specimen.

Inventions 376-420: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus pyogenes in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 645-686, 800, 856, 928 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus pyogenes in a sample or in a clinical specimen.

A kit for the detection of Streptococcus pyogenes in a sample or clinical specimen.

Inventions 421-477: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Klebsiella pneumoniae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 399-448, 792, 794, 829, 899, 902, 903, 934 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Klebsiella pneumoniae in a sample or in a clinical specimen.

A kit for the detection of Klebsiella pneumoniae in a sample or clinical specimen.

Inventions 478-504: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Klebsiella oxytoca in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 449-469, 789, 799, 816, 822, 898, 943 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Klebsiella oxytoca in a sample or in a clinical specimen. A kit for the detection of Klebsiella oxytoca in a sample or clinical specimen.

Inventions 505-571: claims 1-4, 6, 11-12, 13-25 (partially)

An analytical device for direct identification and characterisation of Pseudomonas aeruginosa in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 470-522, 785, 787, 791, 797, 804, 821, 832, 838, 841, 842, 884, 889, 905, 926 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Pseudomonas aeruginosa in a sample or in a clinical specimen.

A kit for the detection of Pseudomonas aeruginosa in a sample or clinical specimen.

Inventions 572-611: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus agalactiae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 606-644, 930 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus agalactiae in a sample or in a clinical specimen.

A kit for the detection of Streptococcus agalactiae in a sample or clinical specimen.

Invention 612: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus mutans in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEQ ID N° 894 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus mutans in a sample or in a clinical specimen.

A kit for the detection of Streptococcus mutans in a sample or clinical specimen.

Inventions 613-633: claims 1-4, 6, 8, 10-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus faecalis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 308-398, 809, 811, 835, 864, 865, 880, 891, 909, 933, 936 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus faecalis in a sample or in a clinical specimen.

A kit for the detection of Enterococcus faecalis in a sample or clinical specimen.

Inventions 634-659: claims 1-4, 6, 8, 10-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus faecuim in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^{\circ}$ 810, 817, 824, 847, 853, 861, 866-874, 876-879, 882, 900, 927, 931, 932, 939, 2887 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus faecuim in a sample or in a clinical specimen.

A kit for the detection of Enterococcus faecuim in a sample or clinical specimen.

Inventions 660-736: claims 1-4, 6, 11-25 (partially)

An analytical device for direct identification and characterisation of Proteus mirabilis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 706-775, 788, 830, 863, 883, 890, 892, 940 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Proteus mirabilis in a sample or in a clinical specimen. A kit for the detection of Proteus mirabilis in a sample or clinical specimen.

Inventions 737-749: claims 1-4, 6, 11-25 (partially)

An analytical device for direct identification and characterisation of Proteus vulgaris in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 776-784, 819, 823, 893, 941 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Proteus vulgaris in a sample or in a clinical specimen. A kit for the detection of Proteus vulgaris in a sample or clinical specimen.

Inventions 750-835: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Candida albicans in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^{\circ}$ 231-307, 910-918 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Candida albicans in a sample or in a clinical specimen. A kit for the detection of Candida albicans in a sample or clinical specimen.

Inventions 836-864: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Acinetobacter baumanii in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 2843-2863, 2865, 2866, 2868-2870, 2888, 2907, 2908 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Acinetobacter baumanii in a sample or in a clinical specimen.

A kit for the detection of Acinetobacter baumanii in a sample or clinical specimen.

Inventions 865-883: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus viridans in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 687-705 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus viridans in a sample or in a clinical specimen.

A kit for the detection of Streptococcus viridans in a sample or clinical specimen.

Invention 884: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Salmonella typhimurium in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEQ ID N° 795 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Salmonella typhimurium in a sample or in a clinical specimen

A kit for the detection of Salmonella typhimurium in a sample or clinical specimen.

Invention 885: claims 1-4, 8, 10-13, 15-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus flavescens in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEQ ID N° 881 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus flavescens in a sample or in a clinical specimen.

A kit for the detection of Enterococcus flavescens in a sample or clinical specimen.

Inventions 886-887: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Staphilococcus hominis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^\circ$ 937, 2906 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Staphilococcus hominis in a sample or in a clinical specimen.

A kit for the detection of Staphilococcus hominis in a sample.

Inventions 888-889: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Stenotrophomonas maltophilia in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 2871, 2875, 2889-2901 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Stenotrophomonas maltophilia in a sample or in a clinical specimen.

A kit for the detection of Stenotrophomonas maltophilia in a sample or clinical specimen.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No
PCT/EP2006/010132

	itent document I in search report		Publication date		Patent family member(s)		Publication date
ΕP	1310569	Α	14-05-2003	CA	2411537		09-05-2003
				JP	2003144153	Α	20-05-2003
				US	2003091991	A1	15-05-2003
WO	9207096	Α	30-04-1992	AT.	161893	T	15-01-1998
				DE	69128639	D1	12-02-1998
				DE		T2	23-04-1998
				DK	554355		11-05-1998
				EP	0554355		11-08-1993
				ES	2112868		16-04-1998
				GR	3026488		31-07-1998
				HK	1005488		08-01-1999
				JP	6502305	 	17-03-1994
US	6747137	B1	08-06-2004	US	2007027309	A1	01-02-2007
ΕP	1344833	Α	17-09-2003	US	2003175716	A1	18-09-2003
US	6008341	A1		NONE			
WO.	02094868		28-11-2002	CA	2440368	A1	 28-11-2002
	1211.300	••	20 21 2002	EP	1373310		02-01-2004
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- (74) Agent: HELBING, Jörg; P.O. Box 10 22 41, 50462 Köln (DE).

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INTERNATIONAL SEARCH REPORT

International application No PCT/EP2006/010132

A. CLASSI	FICATION (OF SUBJECT	MATTER
INV.	C1201/	of subject 68	

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols) C12Q

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, EMBASE, MEDLINE, PAJ, WPI Data, Sequence Search

. DOCUM	ENTS CONSIDERED TO BE RELEVANT		
ategory*	Citation of document, with indication, where appropriate, of	the relevant passages	Relevant to claim No.
4	NAKAMURA M ET AL: "DEVELOPMEN MICRO ARRAY FOR IDENTIFICATIO INFECTIOUS DISEASE CAUSACTIVE HUMAN"	ON OF E BACTERIA IN	1-8, 10-25
	18 May 2003 (2003-05-18), AE THE GENERAL MEETING OF THE AN SOCIETY FOR MICROBIOLOGY, THE WASHINGTON, DC, US, PAGE(S) A XP008047725	MERICAN E SOCIETY,	
	ISSN: 1060-2011 abstract	•	
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X Furth	ner documents are listed in the continuation of Box C.	X See patent family annex.	
Special ca	ategories of cited documents :	"T" later document published after the inte	rnational filing date
A" docume	ont defining the general state of the art which is not ered to be of particular relevance	or priority date and not in conflict with cited to understand the principle or th invention	the application but
	locument but published on or after the international	"X" document of particular relevance; the cannot be considered novel or canno	
L" docume	nt which may throw doubts on priority claim(s) or is cited to establish the publication date of another	involve an inventive step when the do	cument is taken alone
citation	or other special reason (as specified)	"Y" document of particular relevance; the c cannot be considered to involve an in	ventive step when the
other n		document is combined with one or mo ments, such combination being obvio in the art.	
	ent published prior to the international filing date but an the priority date claimed	"&" document member of the same patent	family
ate of the a	actual completion of the international search	Date of mailing of the international sea	
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INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2006/010132

C(Continu		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WANG R-F ET AL: "DNA microarray analysis of predominant human intestinal bacteria in fecal samples" August 2004 (2004-08), MOLECULAR AND CELLULAR PROBES, ACADEMIC PRESS, LONDON, GB, PAGE(S) 223-234, XP004522575 ISSN: 0890-8508 abstract; tables 1,2	1-8, 10-25
A	LEHNER A ET AL: "Oligonucleotide microarray for identification of Enterococcus species" 1 May 2005 (2005-05-01), FEMS MICROBIOLOGY LETTERS, AMSTERDAM, NL, PAGE(S) 133-142, XP004876200 ISSN: 0378-1097 abstract	1-8, 10-25
X	EP 1 310 569 A (PRESIDENT OF GIFU UNIVERSITY) 14 May 2003 (2003-05-14)	1-6, 10-13, 15,19-25
x	WO 92/07096 A (MICROPROBE CORPORATION) 30 April 1992 (1992-04-30) page 12, paragraph 2 page 27, paragraph 2 example 6	1-6,10, 12,13
x	US 6 747 137 B1 (WEINSTOCK KEITH G [US] ET AL) 8 June 2004 (2004-06-08) column 2, lines 41-47 column 16, lines 55-60 column 19, lines 43-61 column 42, lines 5-43 table 2 claim 7	1-6,10, 12,13, 19-25
X	EP 1 344 833 A (CHIP BIOTECHNOLOGY INC DR [TW]) 17 September 2003 (2003-09-17)	1-6, 10-13, 15,16, 18-25
Y	page 2, paragraphs 8,10 page 3, paragraphs 13,18,19 examples 1,2 claim 8	7,14,17
Υ	US 6 008 341 A1 (FOSTER TIMOTHY JAMES [IE] ET AL) 28 December 1999 (1999-12-28) figure 2	7,14,17
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INTERNATIONAL SEARCH REPORT

International application No
PCT/EP2006/010132

ategory*	ion). DOCUMENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
(WO 02/094868 A (CHIRON SPA [IT]; MASIGNANI VEGA [IT]; MORA MARIROSA [IT]; SCARSELLI MA) 28 November 2002 (2002-11-28) page 2, lines 12,13 page 2, lines 20-25 sequences 1992,3983	1-8, 10-15
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International application No. PCT/EP2006/010132

INTERNATIONAL SEARCH REPORT

Box No. II	Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)
This intern	ational search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
	laims Nos.: ecause they relate to subject matter not required to be searched by this Authority, namely:
b	laims Nos.: ecause they relate to parts of the international application that do not comply with the prescribed requirements to such n extent that no meaningful international search can be carried out, specifically:
	laims Nos.: ecause they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box No. III	
This laters	tional Coarching Authority found multiple inventions in this international application, as follows:
inis intern	ational Searching Authority found multiple inventions in this international application, as follows:
	ee additional sheet
	s all required additional search fees were timely paid by the applicant, this international search report covers allsearchable alms.
	s all searchable claims could be searched without effort justifying an additional fees, this Authority did not invite payment of Editional fees.
3. X A	s only some of the required additional search fees were timely paid by the applicant, this international search reportcovers by those claims for which fees were paid, specifically claims Nos.:
	-4 (totally), 5-8, 10-18 (partially), 19-25
	(0000,13), 0 0, 10 10 (000,000)
4. N	o required additional search fees were timely paid by the applicant. Consequently, this international search report is stricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark or	The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.
	The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
	No protest accompanied the payment of additional search fees.
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This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

Invention 1: claims 1-7 and 10-25 (partially)

An analytical device for direct identification and characterisation of microoragnisms in a sample or clinical specimen, wherein the device comprises species specific gene probes of at least 100 nucleotides, and in particular a device for Staphylococcus species identification, in particular for S. aureus identification, wherein, in this case, the microarray comprises the gene probe listed as SEQ ID N° 3 and having a length of at least 100 nucleotides. Use of the analytical device. An in vitro method for identification and characterisation of Staphylococcus aureus in a sample or in a clinical specimen. A kit for the detection of Staphylococcus aureus in a sample or clinical specimen.

Inventions 2-176: claims 1-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus aureus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° SEQ ID N° 1-2, 4-141, 790, 798, 801, 802, 808, 812, 814, 818, 825, 827, 837, 840, 843, 844, 846, 848-852, 854, 855, 859, 862, 875, 885, 896, 897, 904, 907, 908, 935, 942, 2902, 2903, and having a length of at least 100 nucleotides. Use of the analytical device. An in vitro method for identification and characterisation of Staphylococcus aureus in a sample or in a clinical specimen. A kit for the detection of Staphylococcus aureus in a sample or clinical specimen.

Inventions 177-220: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of E. coli in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 142-173, 815, 833, 834, 836, 839, 857, 860, 886-887, 895, 901, 906, and having a length of at least 100 nucleotides. Use of the analytical device. An in vitro method for identification and characterisation of E. coli in a sample or in a clinical specimen.

Inventions 221-258: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus epidermis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 174-208,786, 806, 826 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus epidermis in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus epidermis in a sample or clinical specimen.

Inventions 259-269: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus haemoliyticus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 209-215, 796, 803, 820, 938 and having a length of at least 100 nucleotides. Use of the analytical device.
An in vitro method for identification and characterisation of Staphylococcus haemoliyticus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus haemoliyticus in a sample or clinical specimen.

Inventions 270-276: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus lugdunensis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^\circ$ 216-221, 888 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus lugdunensis in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus lugdunensis in a sample or clinical specimen.

Inventions 277-284: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus warneri in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 224-230, 831 and having a length of at least 100 nucleotides.

Use of the analytical device. An in vitro method for identification and characterisation of Staphylococcus warneri in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus warneri in a sample or clinical specimen.

Inventions 285-286: claims 1-6, 8-25 (partially)

An analytical device for direct identification and characterisation of Staphylococcus saprophyticus in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 222-223 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Staphylococcus saprophyticus in a sample or in a clinical specimen.

A kit for the detection of Staphylococcus saprophyticus in a sample or clinical specimen.

Inventions 287-375: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus pneumoniae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N°523-605, 793, 805, 807, 813, 858, 929 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus pneumoniae in a sample or in a clinical specimen.

A kit for the detection of Streptococcus pneumoniae in a sample or clinical specimen.

Inventions 376-420: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus pyogenes in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 645-686, 800, 856, 928 and having a length of at least 100 nucleotides. Use of the analytical device. An in vitro method for identification and characterisation of Streptococcus pyogenes in a sample or in a clinical specimen. A kit for the detection of Streptococcus pyogenes in a sample or clinical specimen.

Inventions 421-477: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Klebsiella pneumoniae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 399-448, 792, 794, 829, 899, 902, 903, 934 and having a length of at least 100 nucleotides.
Use of the analytical device.

An in vitro method for identification and characterisation of Klebsiella pneumoniae in a sample or in a clinical specimen.

A kit for the detection of Klebsiella pneumoniae in a sample or clinical specimen.

Inventions 478-504: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Klebsiella oxytoca in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 449-469, 789, 799, 816, 822, 898, 943 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Klebsiella oxytoca in a sample or in a clinical specimen. A kit for the detection of Klebsiella oxytoca in a sample or clinical specimen.

Inventions 505-571: claims 1-4, 6, 11-12, 13-25 (partially)

An analytical device for direct identification and characterisation of Pseudomonas aeruginosa in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 470-522, 785, 787, 791, 797, 804, 821, 832, 838, 841, 842, 884, 889, 905, 926 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Pseudomonas aeruginosa in a sample or in a clinical specimen.

A kit for the detection of Pseudomonas aeruginosa in a sample or clinical specimen.

Inventions 572-611: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus agalactiae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 606-644, 930 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus agalactiae in a sample or in a clinical specimen.

A kit for the detection of Streptococcus agalactiae in a sample or clinical specimen.

Invention 612: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus mutans in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEQ ID N $^\circ$ 894 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus mutans in a sample or in a clinical specimen.

A kit for the detection of Streptococcus mutans in a sample or clinical specimen.

Inventions 613-633: claims 1-4, 6, 8, 10-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus faecalis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 308-398, 809, 811, 835, 864, 865, 880, 891, 909, 933, 936 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus faecalis in a sample or in a clinical specimen.

A kit for the detection of Enterococcus faecalis in a sample or clinical specimen.

Inventions 634-659: claims 1-4, 6, 8, 10-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus faecuim in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 810, 817, 824, 847, 853, 861, 866-874, 876-879, 882, 900, 927, 931, 932, 939, 2887 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus faecuim in a sample or in a clinical specimen.

A kit for the detection of Enterococcus faecuim in a sample or clinical specimen.

Inventions 660-736: claims 1-4, 6, 11-25 (partially)

An analytical device for direct identification and characterisation of Proteus mirabilis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 706-775, 788, 830, 863, 883, 890, 892, 940 and having a length of at least 100 nucleotides.

Use of the analytical device. An in vitro method for identification and characterisation of Proteus mirabilis in a sample or in a clinical specimen. A kit for the detection of Proteus mirabilis in a sample or clinical specimen.

Inventions 737-749: claims 1-4, 6, 11-25 (partially)

An analytical device for direct identification and characterisation of Proteus vulgaris in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 776-784, 819, 823, 893, 941 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Proteus vulgaris in a sample or in a clinical specimen. A kit for the detection of Proteus vulgaris in a sample or clinical specimen.

Inventions 750-835: claims 1-6, 8-25 (partially)

clinical specimen.

An analytical device for direct identification and characterisation of Candida albicans in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 231-307, 910-918 and having a length of at least 100 nucleotides. Use of the analytical device. An in vitro method for identification and characterisation of Candida albicans in a sample or in a clinical specimen. A kit for the detection of Candida albicans in a sample or

Inventions 836-864: claims 1-4, 6, 11-12, 14-25 (partially)

An analytical device for direct identification and characterisation of Acinetobacter baumanii in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 2843-2863, 2865, 2866, 2868-2870, 2888, 2907, 2908 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Acinetobacter baumanii in a sample or in a clinical specimen.

A kit for the detection of Acinetobacter baumanii in a sample or clinical specimen.

Inventions 865-883: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus viridans in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 687-705 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus viridans in a sample or in a clinical specimen.

A kit for the detection of Streptococcus viridans in a sample or clinical specimen.

Invention 884: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Salmonella typhimurium in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEO ID N° 795 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Salmonella typhimurium in a sample or in a clinical specimen.

A kit for the detection of Salmonella typhimurium in a sample or clinical specimen.

Invention 885: claims 1-4, 8, 10-13, 15-25 (partially)

An analytical device for direct identification and characterisation of Enterococcus flavescens in a sample or clinical specimen, wherein the microarray comprises the gene probe listed as SEQ ID N $^\circ$ 881 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Enterococcus flavescens in a sample or in a clinical specimen.

A kit for the detection of Enterococcus flavescens in a sample or clinical specimen.

Inventions 886-887: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Staphilococcus hominis in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 937, 2906 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Staphilococcus hominis in a sample or in a clinical specimen.

A kit for the detection of Staphilococcus hominis in a sample.

Inventions 888-889: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Dictyostelium discoideum in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 945, 947 and having a length of at least 100 nucleotides.

Use of the analytical device.

An in vitro method for identification and characterisation of Dictyostelium discoideum in a sample or in a clinical specimen.

A kit for the detection of Dictyostelium discoideum in a sample or clinical specimen.

Inventions 890-892: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Streptococcus dysgalactiae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 2842, 2904, 2905 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Streptococcus dysgalactiae in a sample or in a clinical specimen.

A kit for the detection of Streptococcus dysgalactiae in a sample or clinical specimen.

Inventions 893-907: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Enterobacter cloacae in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N° 2864, 2967, 2872-2874, 2876-2886 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Enterobacter cloacae in a sample or in a clinical

A kit for the detection of Enterobacter cloacae in a sample or clini

Inventions 908-922: claims 1-4, 11-12, 15-25 (partially)

An analytical device for direct identification and characterisation of Stenotrophomonas maltophilia in a sample or clinical specimen, wherein the microarray comprises one of the gene probes listed as SEQ ID N $^\circ$ 2871, 2875, 2889-2901 and having a length of at least 100 nucleotides. Use of the analytical device.

An in vitro method for identification and characterisation of Stenotrophomonas maltophilia in a sample or in a clinical specimen.

A kit for the detection of Stenotrophomonas maltophilia in a sample or clinical specimen.

INTERNATIONAL SEARCH REPORT

Information on patent family members

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Patent document cited in search report	į	Publication date		Patent family member(s)		Publication date
EP 1310569	A	14-05-2003	CA JP US	2411537 2003144153 2003091991	Α	09-05-2003 20-05-2003 15-05-2003
WO 9207096	A	30-04-1992	AT DE DE DK EP ES GR HK JP	2112868 3026488	D1 T2 T3 A1 T3 T3	15-01-1998 12-02-1998 23-04-1998 11-05-1998 11-08-1993 16-04-1998 31-07-1998 08-01-1999 17-03-1994
US 6747137	B1	08-06-2004	US	2007027309	A1	01-02-2007
EP 1344833	Α	17-09-2003	US	2003175716	A1	18-09-2003
US 6008341	A1		NONE			
WO 02094868	Α	28-11-2002	CA EP JP US	2440368 1373310 2005502326 2006115490	A2 T	28-11-2002 02-01-2004 27-01-2005 01-06-2006